SEQUENCE LISTING

<110> The Government of the United States of America, et al.

<120> Recombinant Multivalent Malarial Vaccine Against *Plasmodium Falciparum*

<130> 03063-0440WP

<140>
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<160> 26

<170> PatentIn Ver. 2.0

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<212> DNA
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tct tac atc tat gcg gat cat cat cat cat cat cat aaa cat aaa aaa    96
Ser Tyr Ile Tyr Ala Asp His His His His His His Lys His Lys Lys
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tta aag caa cca ggg gat ggt aat cct tgg tcc cca tgt agt gta act   144
Leu Lys Gln Pro Gly Asp Gly Asn Pro Trp Ser Pro Cys Ser Val Thr
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tgt gga aaa cct aaa gac gaa tta gat tat gaa aat gat att gaa aaa   192
Cys Gly Lys Pro Lys Asp Glu Leu Asp Tyr Glu Asn Asp Ile Glu Lys
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aaa att tgt aaa atg gaa aaa tgt tcc agt gtg ttt aat gtc gta aat   240
Lys Ile Cys Lys Met Glu Lys Cys Ser Ser Val Phe Asn Val Val Asn
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agt aat tct gga tgt ttc aga cat tta gat gaa aga gaa gaa tgt aaa   288
Ser Asn Ser Gly Cys Phe Arg His Leu Asp Glu Arg Glu Glu Cys Lys
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Cys Leu Leu Glu Asp Ser Gly Ser Asn Gly Lys Lys Ile Thr Cys Glu
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tgt tta tta gaa gat tca ggt agc aac gga aag aaa atc aca tgt gaa   336
Cys Leu Leu Glu Asp Ser Gly Ser Asn Gly Lys Lys Ile Thr Cys Glu
            100                 105                 110

tgt act aaa cct gat tct aag cct att gtg caa tat gac aat ttc aat   384
Cys Thr Lys Pro Asp Ser Lys Pro Ile Val Gln Tyr Asp Asn Phe Asn
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gca aac cca aac gca aac ccc aat gca aat cct gat gga aat tgt gaa   432
Ala Asn Pro Asn Ala Asn Pro Asn Ala Asn Pro Asp Gly Asn Cys Glu
    130                 135                 140

gat ata cca cat gta aat gaa ttt tca gca att gat ctt gga aat gct   480
Asp Ile Pro His Val Asn Glu Phe Ser Ala Ile Asp Leu Gly Asn Ala
145                 150                 155                 160

gaa aaa tat gat aaa atg gat gaa cca caa cat tat ggg aaa tca ctc   528
Glu Lys Tyr Asp Lys Met Asp Glu Pro Gln His Tyr Gly Lys Ser Leu
                165                 170                 175

act cca tta gaa gaa tta tat aaa cca aat gat aaa agt ttg tat cag   576
Thr Pro Leu Glu Glu Leu Tyr Lys Pro Asn Asp Lys Ser Leu Tyr Gln
            180                 185                 190

tat ata aaa gca aat tct aaa ttt ata ggt ata act gaa cta agc aac   624
Tyr Ile Lys Ala Asn Ser Lys Phe Ile Gly Ile Thr Glu Leu Ser Asn
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aca ttc ata aac aat gct gga caa cat gga cat atg cat ggt aac gag   672
Thr Phe Ile Asn Asn Ala Gly Gln His Gly His Met His Gly Asn Glu
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agg gaa gat gag aga acg ctt act aag gaa tat gaa gat att gtt ttg   720
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225                 230                 235                 240

aaa gag ttt aca tat atg ata aac ttt gga aga gga cag aat tat tgg   768
Lys Glu Phe Thr Tyr Met Ile Asn Phe Gly Arg Gly Gln Asn Tyr Trp
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gaa cat cca tat caa aaa agt gat caa cct aaa caa tat gaa caa cat   816
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            260                 265                 270

tta aca gat tat gaa aaa att aaa gaa ggt aag ccc ttg gat aaa ttt   864
Leu Thr Asp Tyr Glu Lys Ile Lys Glu Gly Lys Pro Leu Asp Lys Phe
        275                 280                 285

gga aat atc tat gat tat cac tat gag cat tct agt cca tct agt aca   912
Gly Asn Ile Tyr Asp Tyr His Tyr Glu His Ser Ser Pro Ser Ser Thr
    290                 295                 300

aag tca tca agt cca tca aat gta aaa tca gct agt cta gct aca aga   960
Lys Ser Ser Ser Pro Ser Asn Val Lys Ser Ala Ser Leu Ala Thr Arg
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Cys Gly Lys Pro Lys Asp Glu Leu Asp Tyr Glu Asn Asp Ile Glu Lys
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Lys Ile Cys Lys Met Glu Lys Cys Ser Ser Val Phe Asn Val Val Asn
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Ser Asn Ser Gly Cys Phe Arg His Leu Asp Glu Arg Glu Glu Cys Lys
85 90 95

Cys Leu Leu Glu Asp Ser Gly Ser Asn Gly Lys Lys Ile Thr Cys Glu
100 105 110

Cys Thr Lys Pro Asp Ser Lys Pro Ile Val Gln Tyr Asp Asn Phe Asn
115 120 125

Ala Asn Pro Asn Ala Asn Pro Asn Ala Asn Pro Asp Gly Asn Cys Glu
130 135 140

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145 150 155 160

Glu Lys Tyr Asp Lys Met Asp Glu Pro Gln His Tyr Gly Lys Ser Leu
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Thr Pro Leu Glu Glu Leu Tyr Lys Pro Asn Asp Lys Ser Leu Tyr Gln
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Thr Phe Ile Asn Asn Ala Gly Gln His Gly His Met His Gly Asn Glu
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Arg Glu Asp Glu Arg Thr Leu Thr Lys Glu Tyr Glu Asp Ile Val Leu
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Lys Glu Phe Thr Tyr Met Ile Asn Phe Gly Arg Gly Gln Asn Tyr Trp
                245                 250                 255

Glu His Pro Tyr Gln Lys Ser Asp Gln Pro Lys Gln Tyr Glu Gln His
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Leu Thr Asp Tyr Glu Lys Ile Lys Glu Gly Lys Pro Leu Asp Lys Phe
        275                 280                 285

Gly Asn Ile Tyr Asp Tyr His Tyr Glu His Ser Ser Pro Ser Ser Thr
    290                 295                 300

Lys Ser Ser Ser Pro Ser Asn Val Lys Ser Ala Ser Leu Ala Thr Arg
305                 310                 315                 320

Leu Met Lys Lys Phe Lys Ala Glu Ile Arg Asp Phe Phe Gly Ile Ser
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